

### AMENDMENTS TO THE SPECIFICATION

Please delete the existing Sequence Listing and insert the accompanying Sequence Listing (pages 1-118).

Amend the paragraph at page 26, lines 12-18 as follows:

**Table 4.** Illustration of the design of the conformation determining regions and protease binding sites in molecules having P domains larger than 4 amino acids. The P1 residue is underlined. Z is benzyloxycarbonyl group, K[TFA] means Lys(N(epsilon)trifluoroacetyl), Fm is Fmoc (preferably attached to the alpha amino group of the amino terminal residue e.g., e.g., Lysine (K). O indicates tetrahydroisoquinoline-3-carboxylic acid. Aib, designated as B, can be replaced by Pro. J is epsilon-aminocaproic acid (Ahx).~~a-C.~~